In the Sequence Listing

Please enter the attached Sequence Listing, numbered as pages 1-5.

Please substitute the attached Sequence Listing section for the last-filed Sequence Listing.

<110>   KAEMPFER, Raymond
         ARAD, Gila

<120>   BROAD SPECTRUM PYROGENIC EXOTOXINS ANTAGONISTS VACCINES

<130>   KAEMPFER1

<140>   09/150,947
<141>   1998-09-10

<150>   PCT/IL97/00438
<151>   1997-12-30

<150>   ISRAEL 119938
<151>   1996-12-30

<160>   15

<170>   PatentIn version 3.3

<210>   1
<211>   12
<212>   PRT
<213>   Staphylococcus aureus

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1   5   10

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<213>   Staphylococcus aureus


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<221>   LIPID
<222>   (1)..(1)
<223>   N-lauryl cysteine residue

<400>   5

Cys Thr Asn Lys Lys Lys Val Thr Ala Gln Glu Leu Asp
1               5               10


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<220>
<221>   LIPID
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<223>   N-lauryl cysteine residue

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1               5               10


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Tyr Asn Lys Lys Lys Ala Thr Val Gln Glu Leu Asp Tyr Asn Lys Lys
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Lys Ala Thr Val Gln Glu Leu Asp
            20


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<212>   PRT
<213>   Staphylococcus aureus

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Tyr Asn Lys Lys Lys Ala Thr Val Gln Glu Leu Asp Tyr Asn Lys Lys
1               5               10              15


Lys Ala Thr Val Gln Glu Leu Asp Tyr Asn Lys Lys Lys Ala Thr Val
            20              25              30


Gln Glu Leu Asp
            35


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<212> PRT
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<220>
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<222> (14)..(14)
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<220>
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<211> 239
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<213> Staphylococcus aureus

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1           5           10              15
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        20          25              30
```

```
Val Ser Ala Ile Asn Val Lys Ser Ile Asp Gln Phe Leu Tyr Phe Asp
    35          40              45
```

```
Leu Ile Tyr Ser Ile Lys Asp Thr Lys Leu Gly Asn Tyr Asp Asn Val
    50          55              60
```

```
Arg Val Glu Phe Lys Asn Lys Asp Leu Ala Asp Lys Tyr Lys Asp Lys
65          70              75              80
```

```
Tyr Val Asp Val Phe Gly Ala Asn Tyr Tyr Tyr Gln Cys Tyr Phe Ser
            85          90              95
```

```
Lys Lys Thr Asn Asp Ile Asn Ser His Glu Thr Asp Lys Arg Lys Thr
        100         105             110
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```
Cys Met Tyr Gly Gly Val Thr Glu His Asn Gly Asn Gln Leu Asp Lys
        115         120             125
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```
Tyr Arg Ser Ile Thr Val Arg Val Phe Glu Asp Gly Lys Asn Leu Leu
    130         135             140
```

```
Ser Phe Asp Val Gln Thr Asn Lys Lys Lys Val Thr Ala Gln Glu Leu
145         150             155             160
```

```
Asp Tyr Leu Thr Arg His Tyr Leu Val Lys Asn Lys Lys Leu Tyr Glu
            165         170             175
```

```
Phe Asn Asn Ser Pro Tyr Glu Thr Gly Tyr Ile Lys Phe Ile Glu Asn
        180         185             190
```

```
Glu Asn Ser Phe Trp Tyr Asp Met Met Pro Ala Pro Gly Asp Lys Phe
        195         200             205
```

```
Asp Gln Ser Lys Tyr Leu Met Met Tyr Asn Asp Asn Lys Met Val Asp
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```

```
Ser Lys Asp Val Lys Ile Glu Val Tyr Leu Thr Thr Lys Lys Lys
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<210>  13
<211>  10
<212>  PRT
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<220>
<223>  Consensus sequence
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1               5                   10


<210>  14
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<223>  Consensus sequence


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1               5                   10


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<220>
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<223>  Any amino acid

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<223>  Any amino acid

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Xaa Lys Xaa Xaa Xaa Xaa Xaa Xaa Glu Xaa Asp
1               5                   10
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